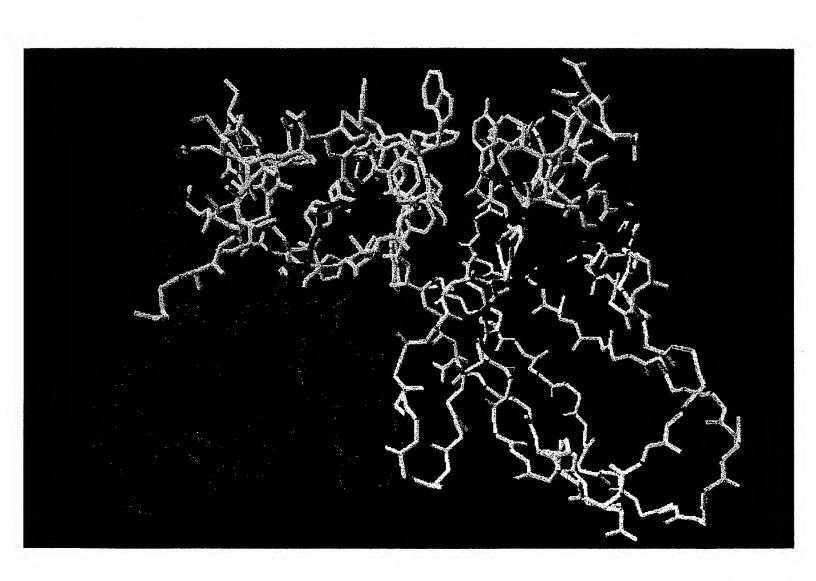
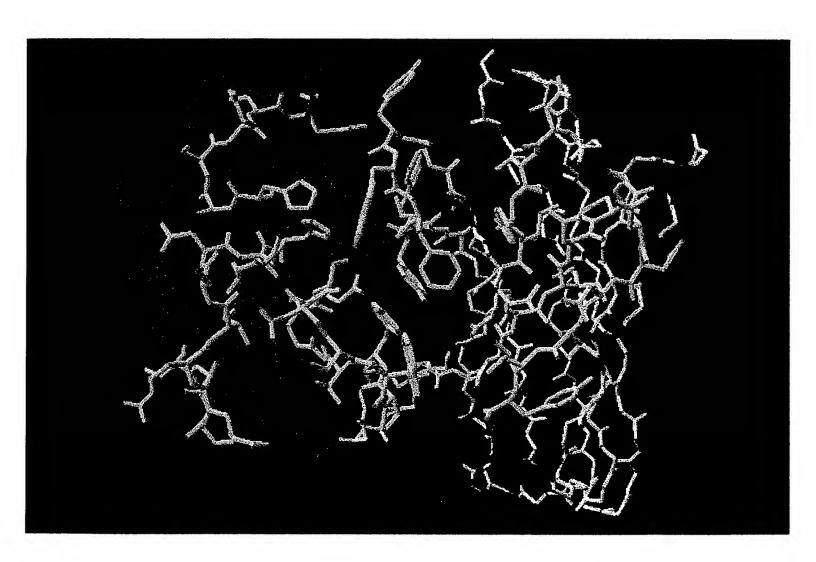
Fig. 2A



)



)

Fig. 2C

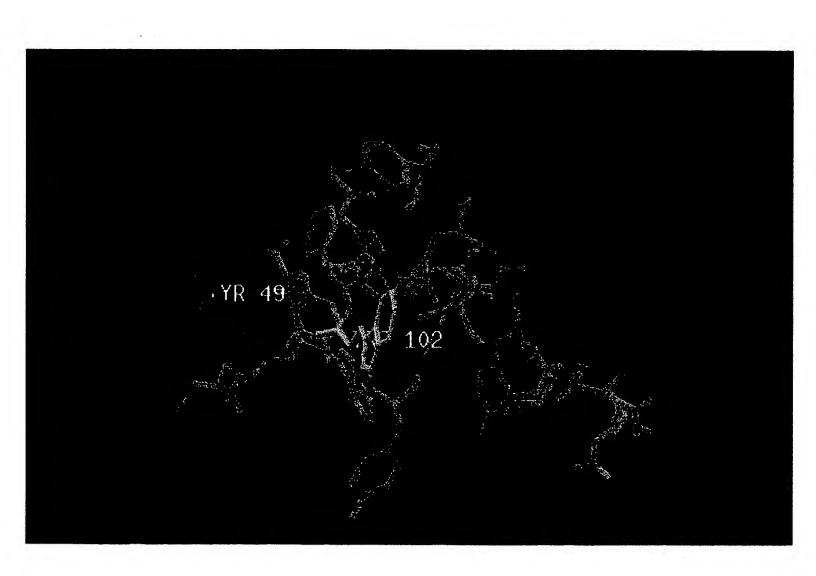


Figure 1. Amino Acid Sequences of Murine 1A6 and Human Consensus Sequences of Heavy Chain Subgroup III (Humiii) and Light Chain κ Subgroup I (Hum κ I).

\mathbf{V}_{H} Domain

Mouse1A6	1 II 21 31 41 EVQLQQSGAE LVKPGASLKL SCTASGFNIK DTYIHWMKQR PEQGLEW I GR ** ** * * * * * * * * * * * * * * * *	
HumIII	EVQLVESGGG LVQPGGSLRL SCAASGFNFSWVRQA PGKGLEWVA—	
Mouse1A6	51 a 61 71 81 a bc 91 IDPANDNTIYD PKVQGKATMT ADTSS NTAYL QLNSLTSEDTAVY YCTT ** ** * * * * * * * * * * * * * * *	
HumIII	A DSVKGRFT IS RDDSKNTAYL QMNSLRAEDTAVY YCTT	
Mouse1A6	103 111 SGYWFA YWGQGTLVT VSS	
HumIII		
$\mathbf{V_L}$ Domain		
Mouse1A6	1 11 21 31 41 51 DIVLTQSPAT LSVTPGDSVS LSCRASQSIS NNLHWYQQKH SESPRLLIKH ASQ ** ** ** ** * ** *	
HumĸI	DIQMTQSPSS LSASVGDRVT ITC WYQQKP GKAPKLLIY	
Mouse1A6	61 71 81 91 101 SISG I PS RFSGSGSGTD FTLSINSVET EDFGMFFCQQ SNSWPYTFGG GTKLEIKR * * * * * * * * * * * * * * * * * * *	
HumĸI	GVPS RFSGSGSGTD FTLTISSLQP EDFATYYC FGQ GTKVEIKR	

The CDR residues as defined by both Kabat and Chothia are shown in boldface.

Figure 3. Amino Acid Sequences of Murine 1A6, Humanized 1A6 (Hum19), and Human Consensus Sequences of Heavy Chain Subgroup III (Humiii) and Light Chain κ Subgroup I (Hum κ I).

$\mathbf{V}_{\mathbf{H}}$ Domain

Mouse	1 11 21 31 41 EVQLQQSGAE LVKPGASLKL SCTASGFNIK DTYIHWMKQR PEQGLEWI GR
Hum19	EVQLVESGGG LVQPGGSLRL SCAASGFNIK DTYIHWVRQA PGKGLEWVAR
HumIII	EVQLVESGGG LVQPGGSLRL SCAASGFNFSWVRQA PGKGLEWVA—
Mouse	51 a 61 71 81 a bc 91 IDPANDNTIYD PKVQGKATMT ADTSS NTAYL QL NSLTSEDTAVY YCT T ** ** * * * * * * * * * * * * * *
Hum19	IDPANDNTIYA DSVKG RFT IS SDDSKNTAYL QMNSLRAEDTAVY YCTA * *
HumIII	A DSVKG RFT IS RDDSKNTAYL QMNSLRAEDTAVY YCTR
Mouse	SGYWFA YWGQGTLVT VSS
Hum19	SGYWFA YWGQGTLVT VSS
HumIII	
	$ m V_L$ Domain
	1 11 21 31 41 51
Mouse	DIVLTQSPAT LSVTPGDSVS LSCRASQSIS NNLHWYQQKH SESPRLLIKH ASQ
Hum19	** ** ** *** * *** DIQMTQSPSS LSASVGDRVT ITCRASQSIS NNLHWYQQKP GKAPKLLIYH ASQ
HumκI	DIQMTQSPSS LSASVGDRVT ITC WYQQKP GKAPKLLIY
	61 71 81 91 101
Mouse	SISG I PS RFSGSGSGTD FTLSINSVET EDFGMFFCQQ SNSWPYTFGG GTKLEIKR * * * * * * * * * * * * * * * * * * *
Hum19	SISGVPS RFSGSGSGTD FTLTISSLQP EDFATYYCQQ SNSWPYTFGQ GTKVEIKR
HumĸI	GVPS RFSGSGSGTD FTLTISSLQP EDFATYYC FGQ GTKVEIKR

The CDR residues as defined by both Kabat and Chothia are shown in boldface.

Figure 4. cDNA Sequences of Humanized scFv3 (Hum3) [SEQ ID. 2].

The restriction sites are underlined. CCATGG NCO I SITE; GGATCC BAMH I SITE;

GTTAAC HPA I SITE

CGAACCATGGGCGATATCcagatgACCCAATCTCCGtctagcCTGAGCgccAG
TgttGGTgatCGAGTTaccattactTGCCGCCCAGCCAATCTATCAGTAATAATCTTC
ACTGGTATCAACAAaaaccgggtaaagctCCGaaaCTTCTTATCAAACACGCCTCTCAG
AGCATTAGCGGCgttCCGAGCCGCTTCTCTGGCTCTGGCTCGGGCACGGACTTT
ACCCTTaccATCAGCTCTcttcagccgGAAGACtttGCCaccTATtatTGTCAGCAGTCTAA
TAGCTGGCCGTATACCTTCGGTcaaGGTACCAAGgtcGAGATTAAGCGCGGCGG
TGGCGGTTCTGGTGGCggtggtagcggtggcGGTGGATCCGGTGGCGGTGGCAGCGA
AGTTCAACTTGTTGAGTCTGGTGGCGGTCTGGTTCAGCCGGTGGCTCTCTGC
GCCTGTCTTGCGCAGCAAGCGGTTTCAACATTAAGGACACCTACATCCATTGG
atgAGGCAAGCTCCGGGTAAGGGTCTGGAGTGGGTGGCACGTATCGACCCGGC
AAACGACAACACCATTTACGATCCGAAGGTGCAGGGCCGTTTTACTatgTCTGC
GGACacCTCTAAGAACACCGCGTACCTTCAGATGAACTCTCTGCGTGCCGAGG
ACACCGCCGTCTACTACTGCACGACCTCTTGGTTAACCCC

Figure 5. Protection of HRV15 infection by mouse 1A6 scFv (Ms1) and humanized 1A6 scFv proteins (Hs3, 4, 7, 17, 18, 19 and 21).

